



<110> SRIKANTHA, THYAGARAJAN SOLL, DAVID R.

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<140> 09/424,951

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<150> PCT/US98/11658

<151> 1998-06-05

<150> 60/048,914

<151> 1997-06-06

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<170> PatentIn Ver. 2.1

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40
45

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85 90 95

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TECH CENTER 1600 2900

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Tyr 225 tta	act Thr	Arg	Gln tcg	Thr	Glu 230 gag	caa Gln	Leu	Leu	Pro gaa	Phe 235 gaa	Ser	Ser gat	His gtg		Val 240 aga	720 768
Tyr 225 tta Leu gat	act Thr ttt Phe	Arg gta Val	Gln tcg Ser	Thr act Thr 245	Glu 230 gag Glu ctt	caa Gln cat His	Leu act Thr	Leu caa Gln ata	Pro gaa Glu 250	Phe 235 gaa Glu ata	Ser ctt Leu ata	Ser gat Asp	His gtg Val aga	Tyr ttg Leu	Val 240 aga Arg	
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Tyr 225 tta Leu gat Asp gaa Glu	act Thr ttt Phe gga Gly gat Asp	gta Val att Ile gca Ala 275 ata	tcg ser ata Ile 260 aca Thr	Thr act Thr 245 gaa Glu ttg Leu att	Glu 230 gag Glu ctt Leu act Thr	caa Gln cat His gga Gly gag Glu	act Thr ttg Leu ccg Pro 280	Leu caa Gln ata 11e 265 gtg Val	gaa Glu 250 cct Pro aaa Lys	Phe 235 gaa Glu ata Ile tat Tyr	Ser ctt Leu ata Ile gat Asp	gat Asp gtg Val ata Ile 285	His gtg Val aga Arg 270 att Ile	Tyr ttg Leu 255 aat Asn atg	Val 240 aga Arg att Ile att Ile	768 816

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tcg atc acc Ser Ile Th													1056
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Gly His Leu	gtg gaa gta gtt Val Glu Val Val 900	gag aac gga Glu Asn Gly 905	ctc gag gcg ta Leu Glu Ala Ty: 91	c Glu Ala
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		Thr Glu Lys	att cga caa tg Ile Arg Gln Tr 940	
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Asp Asn Leu Val Asn Gln Lys Leu Ala Val Arg Ile Leu Glu Lys Gln Gly His Leu Val Glu Val Val Glu Asn Gly Leu Glu Ala Tyr Glu Ala Ile Lys Arg Asn Lys Tyr Asp Val Val Leu Met Asp Val Gln Met Pro 920 Val Met Gly Gly Phe Glu Ala Thr Glu Lys Ile Arg Gln Trp Glu Lys Lys Ser Asn Pro Ile Asp Ser Leu Thr Phe Arg Thr Pro Ile Ile Ala 950 Leu Thr Ala His Ala Met Leu Gly Asp Arg Glu Lys Ser Leu Ala Lys Gly Met Asp Asp Tyr Val Ser Lys Pro Leu Lys Pro Lys Leu Leu Met Gln Thr Ile Lys Lys Cys Ile His Asn Ile Asn Gln Leu Lys Glu Leu Ser Arg Asn Ser Arg Gly Ser Asp Phe Ala Lys Lys Met Thr Arg Asn 1015 Thr Pro Gly Arg Thr Thr Arg Gln Gly Ser Asp Glu Gly Ser Val Lys 1025 Asp Met Ile Gly Asp Thr Pro Arg Gln Gly Ser Val Glu Gly Gly 1050 Thr Ser Ser Arg Pro Val Gln Arg Arg Ser Ala Arg Glu Gly Ser Ile 1060 1065 Thr Thr Ile Ser Glu Gln Ile Asp Arg 1075 <210> 5 <211> 23 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Degenerate primer <220> <221> modified base <222> (16) <223> i <220> <221> modified_base <222> (18) <223> i

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